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## py Correction to: Genetic characterization of Addis Bearded Collies

Gershony, Liza C.

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BMC Genomics. 2020 Dec 30;21(1):898

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CORRECTION

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# Correction to: Genetic characterization of Addison's disease in Bearded Collies

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**Correction to: BMC Genomics (2020) 21:833**  
**<https://doi.org/10.1186/s12864-020-07243-0>**

Following publication of the original article [1], it was noted that due to a typesetting error the caption of Fig. 1 was mistakenly captured within the main body of the article.

Furthermore, the layout of the online versions of Tables 1 and 2 have been updated to improve the presentation of the genotype comparisons.

The correct Fig. 1 with the caption has been included in this Correction article, and the original article has been updated.

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## Reference

1. Gershony LC, Belanger JM, Hytönen MK, et al. Genetic characterization of Addison's disease in Bearded Collies. *BMC Genomics*. 2020;21:833. <https://doi.org/10.1186/s12864-020-07243-0>.

The original article can be found online at <https://doi.org/10.1186/s12864-020-07243-0>.

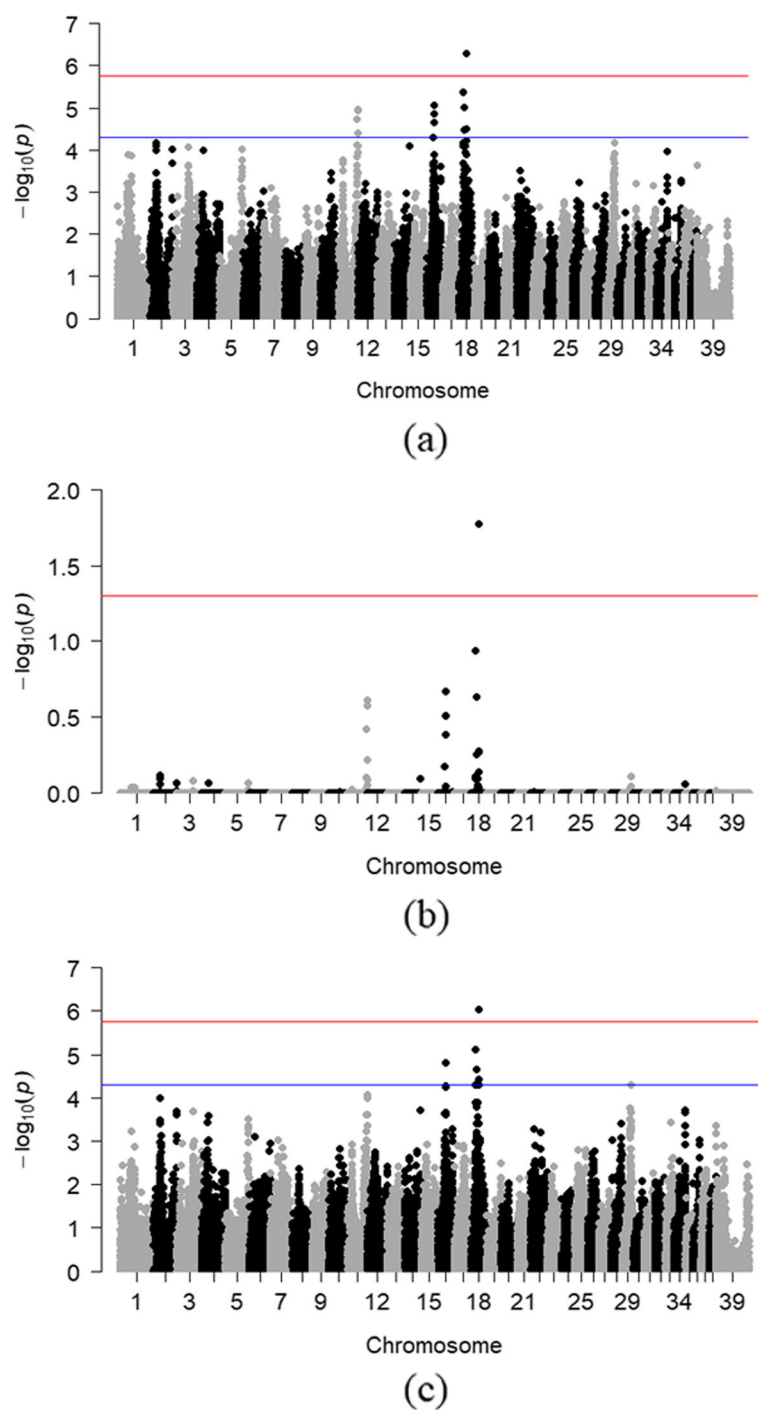
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**Fig. 1** Manhattan Plots. Chi-square based allelic association **(a)** and association testing after 100,000 max(T) permutations **(b)** in PLINK for 103 unrelated Bearded Collies (41 cases, 62 healthy controls;  $\lambda_{GC} = 1.2280$ ). **c** Association testing using GEMMA's univariate linear mixed model approach to account for population substructure of the same dataset ( $\lambda_{GC} = 1.0328$ ). The blue and red lines indicate suggestive ( $p < 0.00005$ ;  $-\log_{10}[p\text{-value}] \geq 4.3$ ) and Bonferroni-adjusted genome-wide significance threshold ( $-\log_{10}[p\text{-value}] \geq 5.75$ ), respectively